



47

1

SEQUENCE LISTING

<110> EMALFARB, MARK A.

PUNT, PETER J.

VAN ZEIJL, CORNELIA

VAN DEN HONDEL, CORNELIUS

<120> HIGH-THROUGHPUT SCREENING OF EXPRESSED DNA LIBRARIES IN  
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<140> 09/834,434

<141> 2001-04-13

<150> PCT/US00/10199

<151> 2000-04-13

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Gly Gln Tyr Ile Leu Gly Asp Thr Val Gly Asp Lys Ile Arg Ile Ile  
 35 40 45

Ala His Tyr Ser Gln Ser Ile Leu Val His Thr Ala Phe Gly Cys Gly  
 50 55 60

Val Leu Thr Ser Ser Thr Arg Met Ser Pro Thr Phe Leu Ser Gln Ser  
 65 70 75 80

Ile Ile Ala Ser Lys Phe Pro Arg Asn Phe Pro Leu Gln Pro Arg Val  
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Tyr Thr Thr Pro Ser Thr Pro Thr Gln Ser Gln Trp Leu Ser Leu Pro  
 100 105 110

Thr Arg Pro Pro Ser Trp Ser Leu Ser Ser Ala Asn Val Leu Thr Phe  
 115 120 125

Gly Thr Phe Thr Leu Lys Ser Gly Arg Arg Ala Ser Pro Leu Gln His  
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Arg His Tyr Arg Asn Arg Lys Thr Tyr His Cys Ile Gln Thr Pro Pro  
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Thr Ser Ser Thr Pro Ala Ser Ser Thr Pro Pro Leu Ser Ser Pro Pro  
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Ser Pro Pro Trp Pro Thr Pro Ser Ser Pro Ser Ser Leu Arg Thr Leu  
 180 185 190

Pro Ser Pro Ser Pro Thr Ser Cys Phe Gly Lys Thr Pro Ser Phe Pro  
 195 200 205

Asn Thr Pro Leu Pro Leu Asn Asn Pro Ile Thr Asn Lys Asn Pro Leu  
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Asn Ser Pro Ala Tyr Lys Gly Ile Pro Leu Ala Cys Ala Thr Leu Leu  
 225 230 235 240

Glu Leu Asn Arg Ile Asp Pro Ala Thr Trp Gly Ser Val Ser Tyr Ser  
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Tyr Asn Arg Lys Glu Ala Lys Asp His Gly Glu Gly Gly Asn Ile Val  
 260 265 270

Gly Ala Ala Leu Lys Gly Lys Thr Val Leu Val Ile Asp Asp Val Ile  
 275 280 285

Thr Ala Gly Thr Ala Met Arg Glu Thr Leu Asn Leu Val Ala Lys Glu  
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Gly Gly Lys Val Val Gly Phe Thr Val Ala Leu Asp Arg Leu Glu Lys  
 305 310 315 320

Met Pro Gly Pro Lys Asp Glu Asn Gly Val Glu Asp Asp Lys Pro Arg  
 325 330 335

Met Ser Ala Met Gly Gln Ile Arg Lys Glu Tyr Gly Val Pro Thr Thr  
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Ser Ile Val Thr Leu Asp Asp Leu Ile Lys Leu Met Gln Ala Lys Gly  
 355 360 365

Asn Glu Ala Asp Met Lys Arg Leu Glu Glu Tyr Arg Ala Lys Tyr Gln  
 370 375 380

Ala Ser Asp Ser Val Ser Leu Thr Asp Cys Leu Gly Gly Cys Glu Arg  
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Leu Gly Val Val Gly Val Gly Met Lys Ser Cys Ile His Arg Gly Leu  
 405 410 415

Lys Arg Cys Val Glu Thr Val Val Arg Cys Phe Met Ser Lys Ser Thr  
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Asn Asp Thr Leu Lys Lys Thr Pro Trp Phe Gln Leu Asn Pro Gly Lys  
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Met Leu Gly Thr Pro Val Pro Thr Gln Trp Ala Pro Val Ser His Ile  
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Ser Gly Arg Arg Leu Phe Gly Gly Cys Gly Leu Glu Arg His Tyr Gly  
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 35 40 45

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His Pro Pro Glu Ala Gln Leu Phe Cys Leu Asn Gln Leu His Pro Asn  
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Ser Pro Ala Thr Ser Pro Ser Asn Pro Val Ser Ile Pro Leu Pro Pro  
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His Pro His Asn His Asn Gly Ser Pro Cys Leu Gln Asp Arg Leu Pro  
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Gly Val Ser Arg Arg Pro Thr Cys Leu Ser Ala Pro Ser Pro Ser Arg  
 115 120 125

Val Ala Val Arg His Pro Ser Asn Thr Gly Ile Ile Ala Ile Gly Arg  
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Leu Thr Thr Val Tyr Arg Leu Pro Leu Leu Leu Gln Arg Arg His Leu  
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Gln His Arg Leu Ser Pro Leu Arg Pro Leu His His Gly Pro His His  
 165 170 175

His His Leu Pro Arg Glu Pro Phe His Pro Gln Ala Arg Arg His Ala  
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Ser Gly Lys Lys Pro Pro Leu Ser Pro Ile Pro His Phe His Ser Thr  
 195 200 205

Thr His Lys Leu Thr Lys Thr Pro Thr Ala Pro His Thr Lys Ala Ser  
 210 215 220

Pro Ser Arg Ala Pro Pro Ser Leu Asn Ser Thr Ala Ser Thr Pro Pro  
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Pro Gly Ala Ala Cys Pro Thr Ala Thr Ala Lys Lys Pro Arg Ile  
 245 250 255

Thr Ala Lys Ala Ala Thr Leu Ser Ala Pro Leu Arg Ala Arg Pro Cys  
 260 265 270

Leu Ser Thr Met Ser Ser Arg Pro Val Pro Pro Cys Val Arg Pro Ser  
 275 280 285

Thr Trp Ser Pro Arg Arg Ala Ala Arg Ser Ser Asp Ser Leu Leu Leu  
 290 295 300

Trp Thr Ala Trp Arg Arg Cys Pro Asp Pro Arg Thr Arg Thr Val Ser  
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 325 330 335

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 Ile Arg Leu Val Ile Ser Arg Phe His Pro Ile Val Trp Val Gly Val  
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 Arg Gly Val Arg Leu Trp Ala Glu Lys Ala Val Tyr Ile Gly Ala Arg  
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 Glu Trp Pro Ile Tyr Pro Arg Arg Tyr Arg Arg Arg Asp Lys Asn Asn  
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 Arg Thr Leu Phe Pro Lys His Thr Gly Thr Tyr Cys Ile Arg Leu Val  
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 Arg Gly Ala Tyr Leu Ile His Pro Asn Glu Pro Asn Phe Phe Val Ser  
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 Ile Asn Asn Cys Ile Gln Ile Pro Pro Gln Leu Pro Pro Pro Thr Pro  
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 Cys Leu Tyr His Ser Leu His Thr His Thr Ile Thr Met Ala Leu Pro  
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Ala Tyr Lys Thr Ala Phe Leu Glu Ser Leu Val Gly Gln Arg Ala Asp  
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 Phe Arg His Leu His Pro Glu Val Gly Ser Pro Cys Val Thr Pro Pro  
 130 135 140  
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 145 150 155 160  
 Pro Tyr Phe Phe Asn Ala Gly Ile Phe Asn Thr Ala Ser Leu Leu Ser  
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 Ala Leu Ser Thr Met Ala His Thr Ile Ile Thr Phe Leu Ala Glu Asn  
 180 185 190  
 Pro Ser Ile Pro Lys Pro Asp Val Met Leu Arg Val Lys Asn Pro Leu  
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 Phe Pro Gln Tyr Pro Thr Ser Thr Gln Gln Pro Ile Asn Asn Gln Lys  
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 Pro Pro Lys Gln Pro Arg Ile Gln Arg His Pro Pro Arg Val Arg His  
 225 230 235 240  
 Pro Pro Thr Gln Pro His Arg Pro Arg His Leu Gly Gln Arg Val Leu  
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 Gln Leu Gln Pro Gln Arg Ser Gln Gly Ser Arg Arg Arg Gln His  
 260 265 270  
 Cys Arg Arg Arg Ser Glu Gly Gln Asp Arg Ala Cys Asp Arg Arg Cys  
 275 280 285  
 His His Gly Arg Tyr Arg His Ala Asp Pro Gln Pro Gly Arg Gln Gly  
 290 295 300  
 Gly Arg Gln Gly Arg Arg Ile His Cys Cys Ser Gly Pro Leu Gly Glu  
 305 310 315 320  
 Asp Ala Arg Thr Gln Gly Arg Glu Arg Cys Arg Gly Arg Ala Gln Asn  
 325 330 335  
 Glu Cys Tyr Gly Ser Asp Pro Gly Val Trp Cys Ala His Asp Glu Tyr  
 340 345 350  
 Cys Tyr Ser Gly Phe Asp Gln Val Asp Ala Gly Glu Gly Gln Gly Arg  
 355 360 365  
 Tyr Glu Ala Val Gly Gly Val Gly Val Ser Gly Leu Val Gly Phe Ile  
 370 375 380  
 Asp Arg Leu Phe Gly Trp Val Glu Val Arg Leu Gly Cys Gly Arg Arg  
 385 390 395 400  
 Asn Glu Lys Leu Tyr Thr Gly Pro Glu Glu Val Arg Arg Asp Gly Arg  
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Glu Met Phe Tyr Val Lys Ile Leu Asn Lys His Leu Lys Lys Asp Pro  
420 425 430

Leu Val Ser Ala Glu Leu Ala Arg Lys Asp Ala Arg His Ala Met Ser  
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Leu Ala His Ser Val Gly Thr Arg Phe Pro His Leu Lys Trp Pro Thr  
450 455 460

Leu Ile Trp Leu Arg Leu Trp Pro Gly Lys Ala Leu Trp Arg Ala Ala  
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Val Gln Gly Arg Gly Trp Arg Thr Asn His Asp Ala Arg Arg Glu Leu  
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